Conservation Genetics and Captive Breeding

Peter Dratch, NPS & Tim King, USGS

Channel Island Fox





Captive Breeding of Channel Island Fox

Robert Wayne and Melissa Gray UCLA Conservation Genetics Resource Center

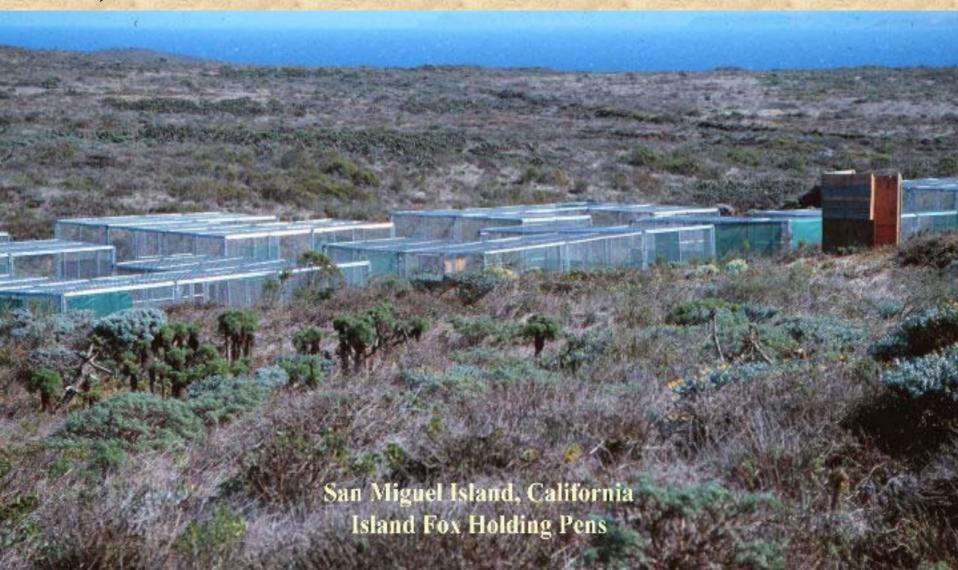
Coleen Lynch AZA Population Management Center

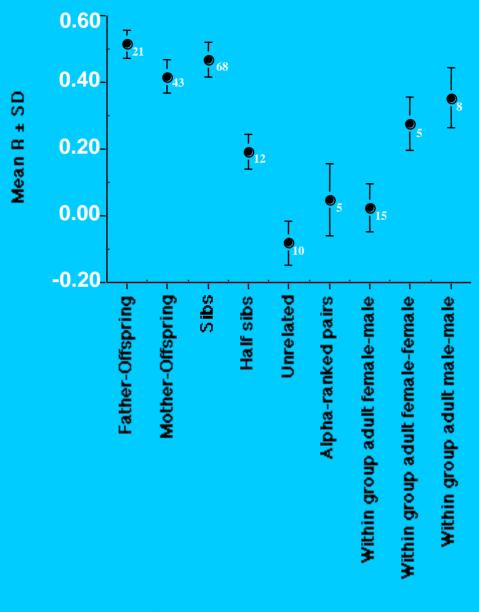
Tim Coonan Channel Islands National Park



Genetic Management

- A) Which individuals should be bred and paired?
- B) Which individuals should be reintroduced?





Social Grouping

Relatedness of Adults

San Miguel

	Sex	M	F	M	F	F	F	M	
Sex	Individual	44829	90D1A	47B06	11929	60921	E2677	85D02	
M	44829	*							
F	90D1A	-0.070	*						
M	47B06	0.160	0.425	*					
F	11929	0.750	0.245	-0.069	*				
F	60921	-0.640	-0.189	-0.405	-0.371	*			
F	E2677	-0.191	0.478	0.058	0.134	0.400	*		
M	85D02	0.505	-0.059	-0.066	0.509	-0.197	-0.217	*	
F	92C32	0.736	0.172	0.375	0.467	-0.467	0.077	0.042	
F	7534A	-0.859	0.117	-0.145	-0.587 0.80		0.013	-0.190	
F	F6558	-0.135	-0.514	-0.772	-0.120	-0.120 0.374		0.192	
M	7574A	-0.514	-0.086	0.525	-0.743	-0.743		-0.516	
M	57150	0.481	0.040	-0.541	0.748	-0.162	0.125	0.699	
F	71071	-0.181	0.597	0.411	-0.167	-0.306	0.425	0.004	
F	B0B25	0.167	-0.302	-0.244	0.176	0.289	0.028	0.386	
F	92804	-0.098	-0.527	0.386	-0.697	0.022	-0.273	-0.198	
F	61B03	-0.493	-0.299	0.108	-0.758	0.459	-0.103	-0.395	

SM r-values < -0.12 SR r-values < 0.08

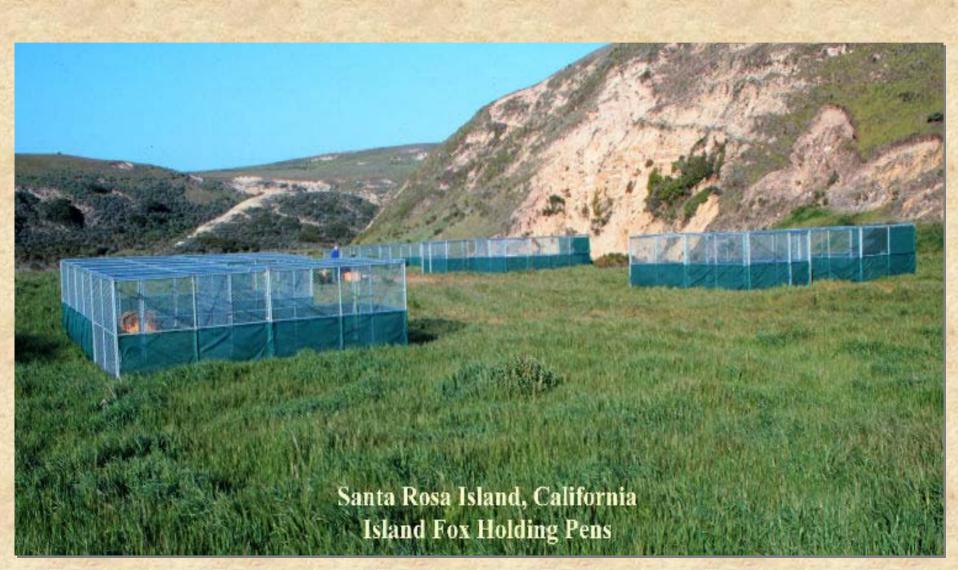
Relatedness of San Miguel pups

	Sai	n Miguel	Pups (M)									
Ì	Jai	ii wiiguei		Parents of	of pups							
i,	Poter	ntial mates	C4A16	/C7303	70C1D/83C24/11F73							
	Sex	ID	92C32 (F)	7574A (M)	90D1A (F)	44829 (M)						
5	M	44829	0.736	-0.514	-0.070	*						
	F	90D1A	0.172	-0.086	*	-0.070						
	M	47B06	0.375	0.525	0.425	0.160						
ñ	F	11929	0.467	-0.743	0.245	0.750						
ľ	F	60921	-0.467	0.233	-0.189	-0.640						
	F	E2677	0.077 -0.181		0.478	-0.191						
ģ	М	85D02	0.042	-0.516	-0.059	0.505						
3	F	92C32	*	-0.349	0.172	0.736						
	F	7534A	-0.668	0.436	0.117	-0.859						
	F	F6558	-0.217 -0.421		-0.514	-0.135						
	M	7574A	-0.349	*	-0.086	-0.514						
ij	M	57150	0.184	-1.024	0.040	0.481						
	F	71071	-0.256	0.071	0.597	-0.181						
5	F	B0B25	0.114	-0.488	-0.302	0.167						
	F	92804	-0.179	0.550	-0.527	-0.098						
	F	61B03	-0.302	0.697	-0.299	-0.493						

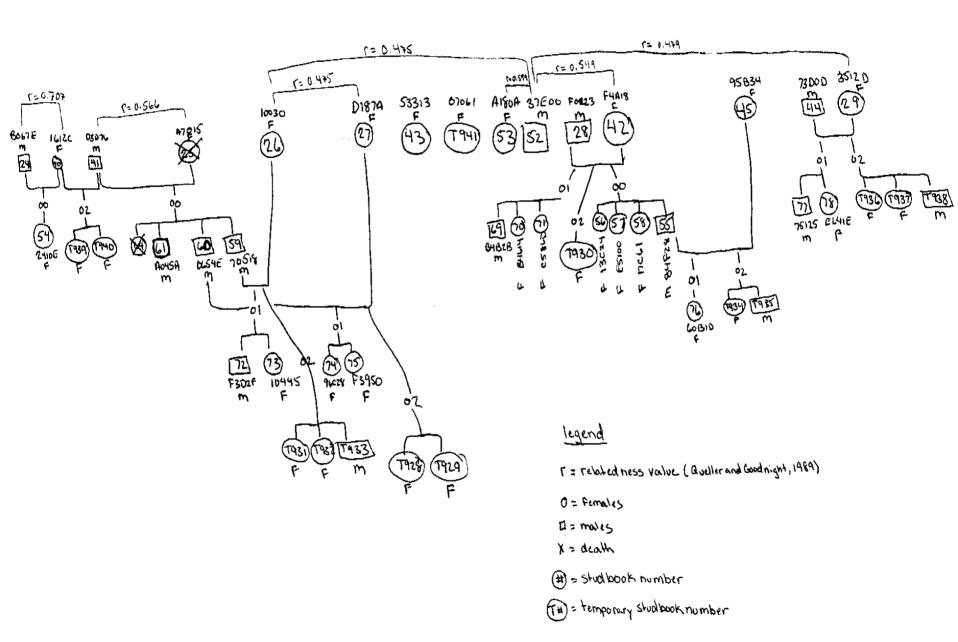


SM r-values < -0.12

Santa Rosa Island 2000: 14 individuals, all but 1 in captivity



channel bland Fox Pedigree: Santa Rosa



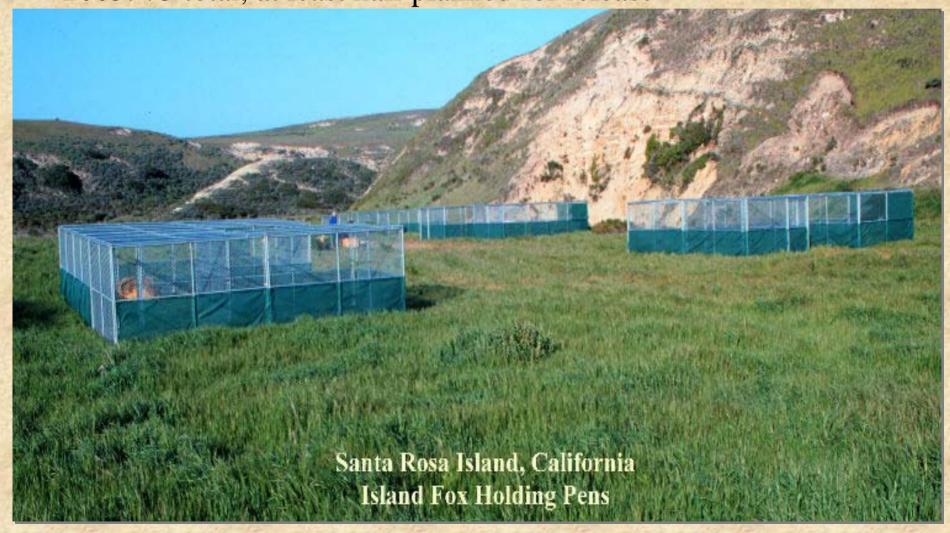
San Mig	uel		Rank	ing			Santa R	losa		Ranl	king	
ID	sex	1ST	2ND	3RD	4TH		ID	sex	1ST	2ND	3RD	4TH
C7303	M	F6558	71071	E2677			34614	F	D3D76	F3D2F		
C4A16	M	F6558	71071	E2678					B067E			
70C1D	M	60921	92804			S Anna (C)	C586D	F	D3D76	F3D2F		
		F6558							B067E			
		61B03					B4B2B	М	2410E	10445		
83C24	M	60921	92804			9 0			1612C			
		F6558					10445	F	B4B2B	A045A		
		61B03					F3D2M	М	E5100	34614		
11F73	М	60921	92804							C586D		
		F6558				THE REPORT OF THE PARTY OF THE	96C2E	F	37E00	75125	B4B2B	
		61B03					F3950	F	37E01	75126	B4B2B	
85D02	M	61B03	92804				60B1D	F	A045A	75125	B067E	
		E2677	60921				75125	М	53313	60B1D		
			7534A				E6D1E	F	A045A			
57150	M	61B03	7534A	71071		Suggested parings	D3D76	М	E5100	37C61	13C24	
		92804		60921		buggested parmgs			34614	53313	A180A	
47B06	М	F6558	60921	7534A	11929	0			C586D		1612C	
			BOB25			& reintroduction	B067E	M	34614	53313	13C24	E5100
11929	F	47B06							C586D		A180A	
71071	F	57150	C7303			for San Miguel					37C61	
			C4A16			101 San Wilguel	37E00	М	96C2E	1612C		
60921	F	47B06	70C1D	85D02		1 C D			F3950	2410E		
			83C24	57150		and Santa Rosa				53313		
			11F73							2410E		
F6558	F	47B06	C7303	70C1D		PARTY NAMED IN COLUMN 2015	A045A	M	53313	13C24	E5100	10445
			C4A16	83C24		TO THE RESIDENCE OF THE PARTY AND THE PARTY			60B1D	A180A	37C61	
				11F73		经济的政策,不是 第二条的基础的经	E5100	F	F3D2F	A045A	B067E	
7534A	F	57150	47B06						D3D76		75125	
			85D02				2410E	F	B4B2B		37E00	
BOB25	F	47B06					13C24	F	A045A	D3D76		
92804	F	57150	85D02	70C1D			A180A	F	D3D76	A045A		
				83C24					B067E			
				11F73			1612C		B4B2B			
E2677	F	85D02	C7303				37C61	F	D3D76	B067E	A045A	
			C4A16						F3D2F			
			47B06				53313	F	F3D2F			
61B03	F	57150	85D02			Island Fox (Urocyon littoralis)			75125			
			70C1D						D3D76			
			83C24						B067E			
			11F73						A045A			

Santa Rosa Island

2000: 14 individuals, all but 1 in captivity

2004: 45 total, 16 males/29 females, first releases

2005: 73 total, at least half planned for release



Island Fox Rescue Depends Critically on Cooperation

NPS FWS NC UCLA **UCDavis USGS** CIC AZA/ **PMC CFG CESU**



Captive Breeding Management (Ex situ conservation)

The Robust Unique Multilocus Genotype



Captive Breeding Management

Captive Breeding Management Plan = Species Survival Plan

Plan must facilitate the maintenance of a genetically viable and demographically stable population of a species in captivity (95 % chance of survival for 100 years (Ballou)).

Establish a metapopulation structure (subpopulations in geographically distinct areas)

Prerequisites and Assumption

Prerequisites
Phylogeographic and phylogenetic relationships within and among close relatives is known

Sufficient genetic diversity to realize unique multilocus genotypes

Assumption
Preserving maximum levels of genetic diversity within and among populations will increase fitness

Captive Breeding Successes

- ·19% of all mammals, 10% of all bird species have been bred in captivity.
- •90% of all mammals, 74% of all birds added to U.S. zoo collections since 1985 were born in captivity.
- •Some species are extinct in the wild but thrive in zoos: Przewalski's horse, Arabian Oryx, Pere David's deer.
- ·A number of wild populations of species were born in captivity and now live free: Bald Eagle, whooping crane, Andean condors, red wolves, Golden Lion Tamarin.
- ·A successful captive breeding program by USFWS with a bobwhite quail generated the creation of a wildlife refuge in southern Arizona to allow its successful reintroduction.
- ·Public awareness and concern can be mobilized by such efforts.

Unique Multilocus Genotypes

Sfo Loci

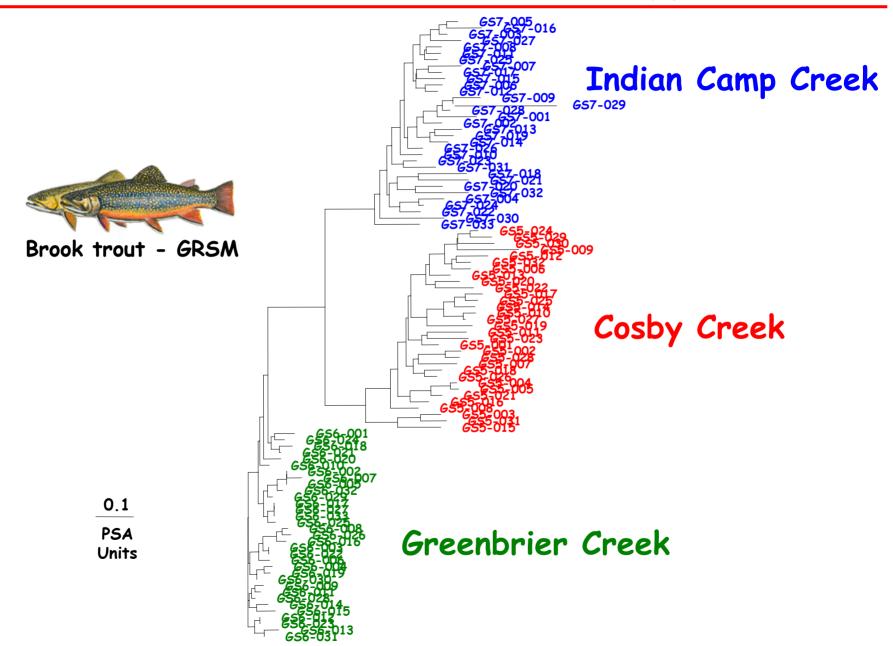
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MD7-18	187	215	116	119	175	183	140	146	123	123	113	116	178	187	139	139	333	341	221	233	208	208	212	240	226	230
MD7-19	201	225	158	158	183	199	143	143	123	123	101	110	181	193	139	157	365	365	227	233	200	212	240	240	214	230
MD7-20	201	215	116	122	175	191	143	146	120	123	110	116	181	187	136	139	341	365	233	233	200	204	212	216	238	250
MD7-21	215	225	110	113	175	183	140	146	123	123	113	122	178	193	139	151	333	337	230	233	212	216	212	244	226	234
MD7-22	225	225	113	119	175	199	143	146	120	123	113	116	181	187	130	157	337	365	227	230	208	216	212	240	214	238

Alleles measured in base pairs



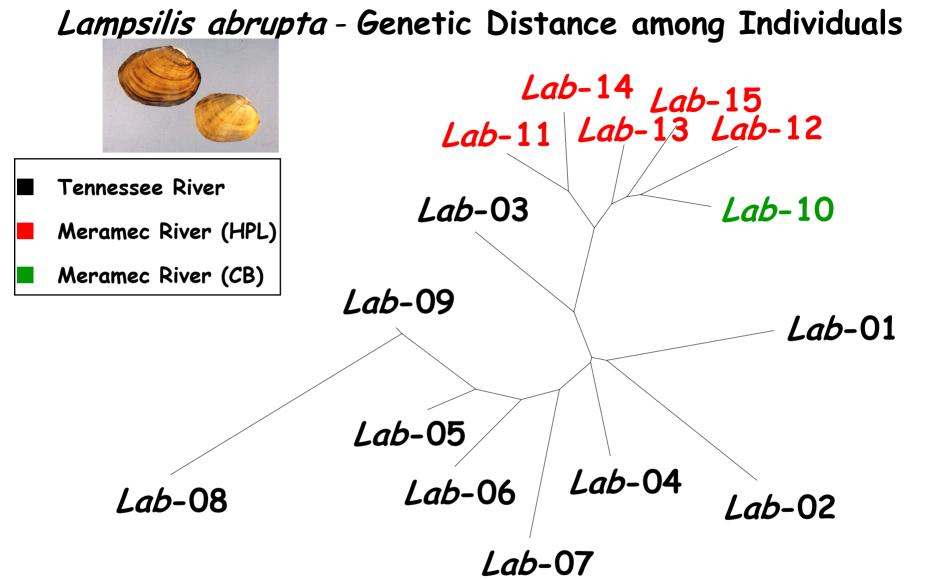
Brook trout - GRSM

Unique Multilocus Genotypes



Applications of Unique Multilocus Genotypes

- ·Delineate fine-scale population structure
- ·Genetic Stock Identification
- ·Enhanced assignment or allocation analysis
- ·Hybridization
- · Movement
- ·Kinship



Neighbor-Joining Tree - Proportion of Shared Alleles Multilocus Genotypes from 15 Microsatellite DNA Loci

0.1

Enlightened Broodstock Management



Assess baseline levels of genetic diversity and variation Determine multi-locus genotypes

Increase levels of heterozygosity Calculate genetic distance (PSA) between individuals Cross distantly related individuals to increase heterozygosity (and alter the distribution of alleles)

Result: heterozygosity increased 6.2% in one generation

Increase levels of genetic diversity Introduce new alleles from other populations (if necessary)



Proportion of Shared Alleles (PSA) Distance



Lampsilis

abrupta

Lampsilis abrupta

Lab-002 Lab-004

Lab-002 Lab-008

Lab-003 Lab-015

Lab-004 Lab-005

Lab-004 Lab-010

Lab-005 Lab-010

1.792

1.792

1.792

1.792

1.792

1.792

Animal 1 Animal 2	PSA	Animal 1	Animal 2	PSA	Animal 1 Animal 2	PSA
Lab-011 Lab-014	0.836	Lab-005	Lab-015	1.792	Lab-002 Lab-011	2.303
Lab-010 Lab-015	0.916	Lab-006	Lab-007	1.792	Lab-003 Lab-005	2.303
Lab-013 Lab-015	0.916	Lab-006	Lab-009	1.792	Lab-003 Lab-008	2.303
Lab-010 Lab-012	1.003	Lab-006	Lab-015	1.792	Lab-003 Lab-012	2.303
Lab-010 Lab-013	1.003	Lab-009	Lab-011	1.792	Lab-006 Lab-011	2.303
Lab-011 Lab-013	1.003	Lab-011	Lab-012	1.792	Lab-007 Lab-014	2.303
Lab-012 Lab-013	1.003	Lab-014	Lab-015	1.792	Lab-007 Lab-015	2.303
Lab-013 Lab-014	1.099	Lab-001	Lab-002	2.015	Lab-009 Lab-015	2.303
Lab-005 Lab-006	1.204	Lab-001	Lab-005	2.015	Lab-001 Lab-003	2.708
Lab-011 Lab-015	1.204	Lab-001	Lab-006	2.015	Lab-001 Lab-010	2.708
Lab-012 Lab-015	1.204	Lab-001	Lab-007	2.015	Lab-002 Lab-007	2.708
Lab-012 Lab-014	1.322	Lab-001	Lab-015	2.015	Lab-002 Lab-010	2.708
Lab-003 Lab-004	1.455	Lab-002	Lab-005	2.015	Lab-002 Lab-012	2.708
Lab-005 Lab-009	1.455	Lab-002	Lab-009	2.015	Lab-002 Lab-013	2.708
Lab-008 Lab-009	1.455	Lab-002	Lab-014	2.015	Lab-004 Lab-011	2.708
Lab-010 Lab-011	1.455	Lab-003	Lab-006	2.015	Lab-005 Lab-013	2.708
Lab-010 Lab-014	1.455	Lab-003	Lab-009	2.015	Lab-005 Lab-014	2.708
Lab-001 Lab-004	1.609	Lab-003	Lab-011	2.015	Lab-006 Lab-012	2.708
Lab-001 Lab-011	1.609	Lab-003	Lab-014	2.015	Lab-006 Lab-013	2.708
Lab-003 Lab-007	1.609	Lab-004	Lab-008	2.015	Lab-006 Lab-014	2.708
Lab-003 Lab-010	1.609	Lab-004	Lab-009	2.015	Lab-007 Lab-008	2.708
Lab-003 Lab-013	1.609	Lab-004	Lab-012	2.015	Lab-009 Lab-010	2.708
Lab-004 Lab-006	1.609	Lab-004	Lab-013	2.015	Lab-009 Lab-014	2.708
Lab-004 Lab-007	1.609	Lab-004	Lab-014	2.015	Lab-001 Lab-012	3.401
Lab-004 Lab-015	1.609	Lab-005	Lab-012	2.015	Lab-002 Lab-015	3.401
Lab-005 Lab-007	1.609	Lab-006	Lab-008	2.015	Lab-005 Lab-011	3.401
Lab-005 Lab-008	1.609	Lab-006	Lab-010	2.015	Lab-007 Lab-010	3.401
Lab-007 Lab-009	1.609	Lab-007	Lab-011	2.015	Lab-007 Lab-013	3.401
Lab-002 Lab-003	1.792	Lab-007	Lab-012	2.015	Lab-008 Lab-011	3.401

Lab-009 Lab-013

Lab-001 Lab-008

Lab-001 Lab-009

Lab-001 Lab-014

Lab-002 Lab-006

Lab-001

Lab-013

2.015

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Lab-008 Lab-012

Lab-008 Lab-013

Lab-008 Lab-014

Lab-009 Lab-012

Lab-008 Lab-010

Lab-008 Lab-015

3.401

3.401

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10

10

The Connecticut River



Experiment

Gene marking - 2 for the price of 1; will determine levels of genetic variability and allow assessment of all aspects of the stocking program

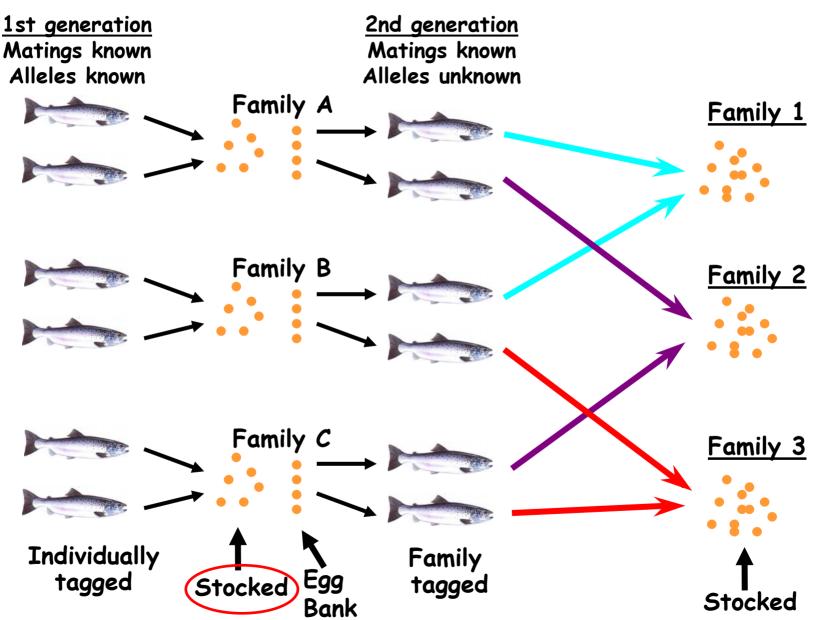
Determine CR tributaries with the greatest production

Determine the favorable characteristics of productive tributaries

Ultimately achieve restoration of a reproducing population

The Connecticut River





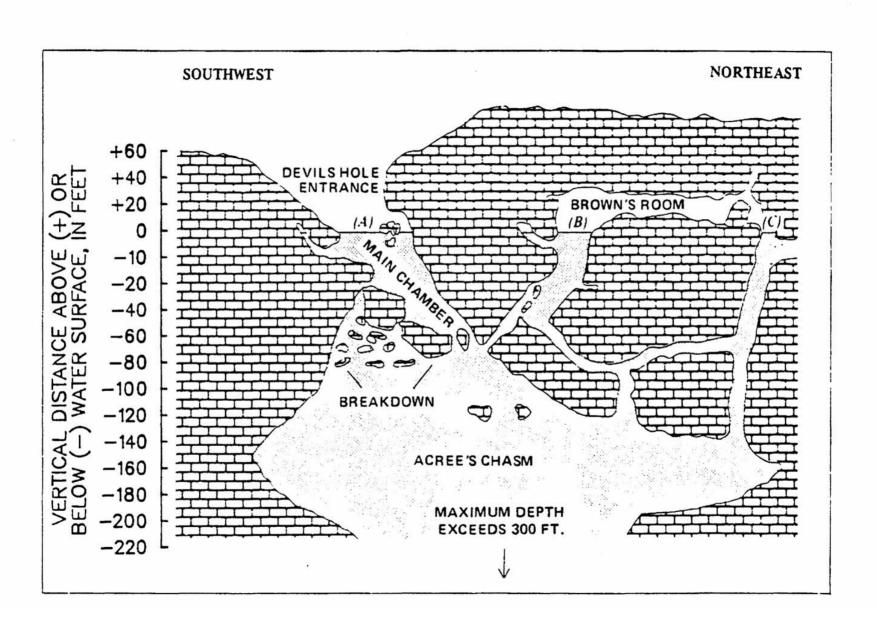
Devils Hole Pupfish (Cyprinodon diabolis)



An Ongoing Lesson in Conservation Biology with thanks to Andrew Martin (U Colorado) and John Wullschleger (NPS Water Resources)





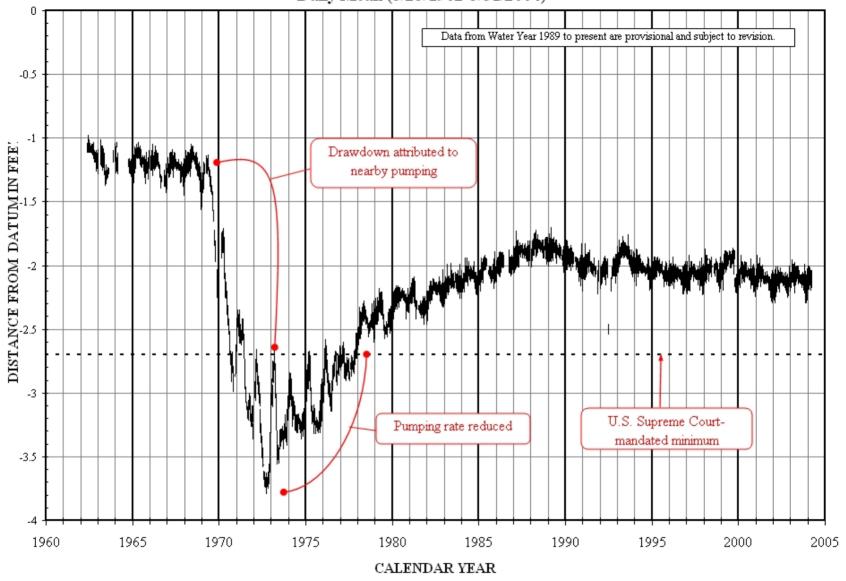


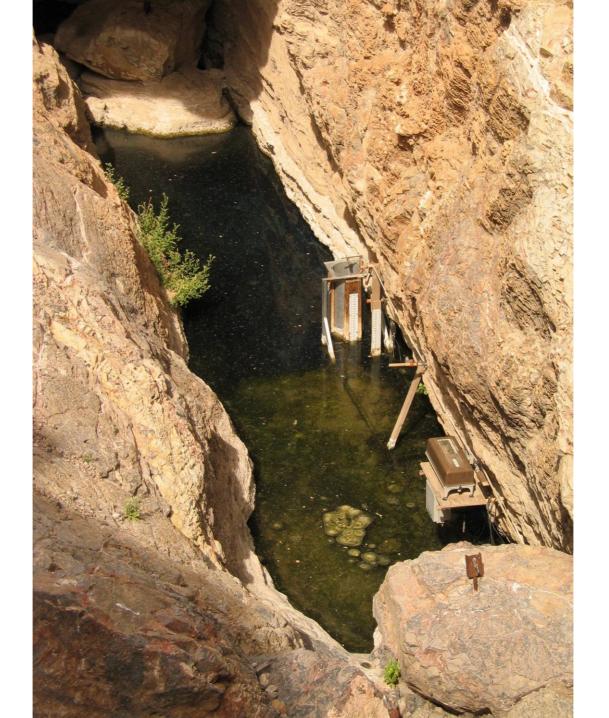
Devils Hole Abbreviated Chronology

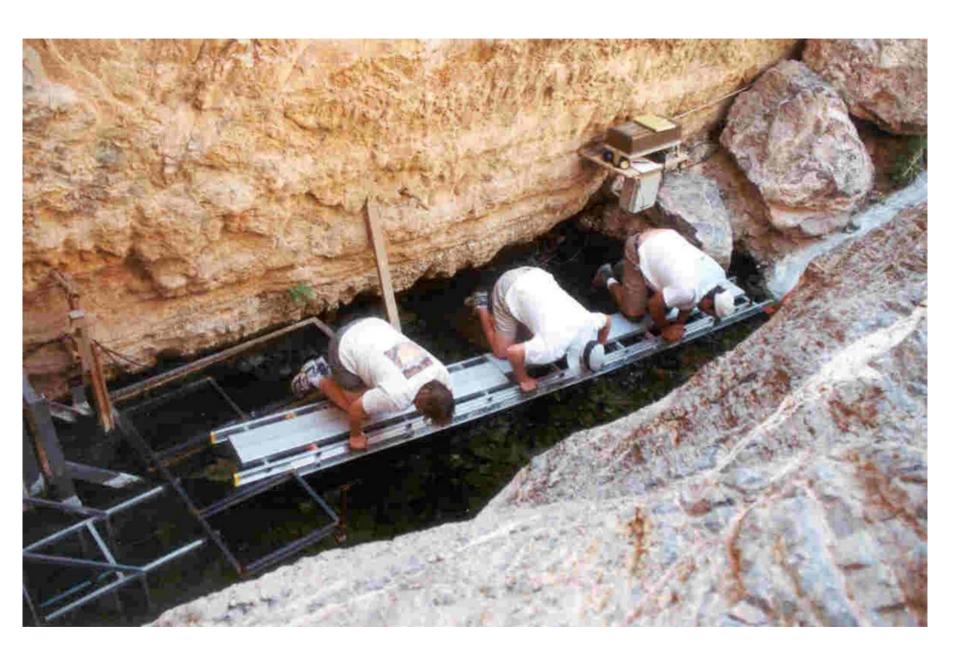
- 1952 Devils Hole is protected as a 40 acre disjunct unit of Death Valley National Monument.
- 1967 Devils Hole pupfish is listed as an endangered species.
- 1976 Supreme Court concludes that when NPS acquired Devils Hole the federal government implicitly reserved sufficient water to protect the pupfish and its habitat.

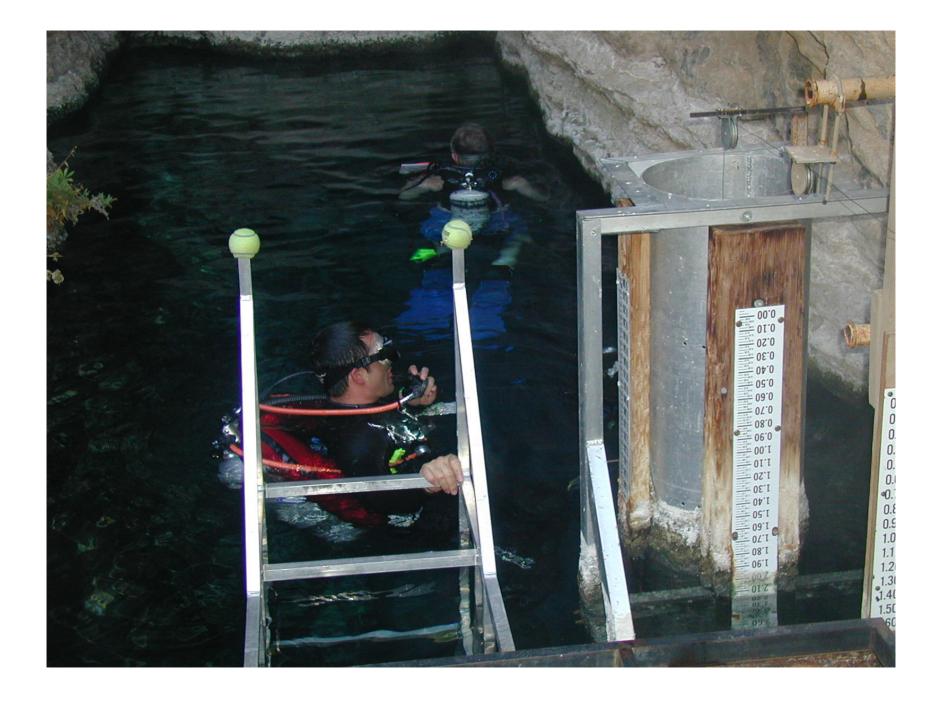
 A lower court then established minimum water level.
- 1984 Ash Meadows National Wildlife Refuge is established, in part to prevent groundwater pumping near to Devils Hole.
- 1994 Death Valley and Devils Hole become a National Park.
- 2004 Flash flood kills at least 80 adult pupfish in Devils Hole.

Pool Stage in Devils Hole Daily Mean (5/23/1962-3/31/2004)

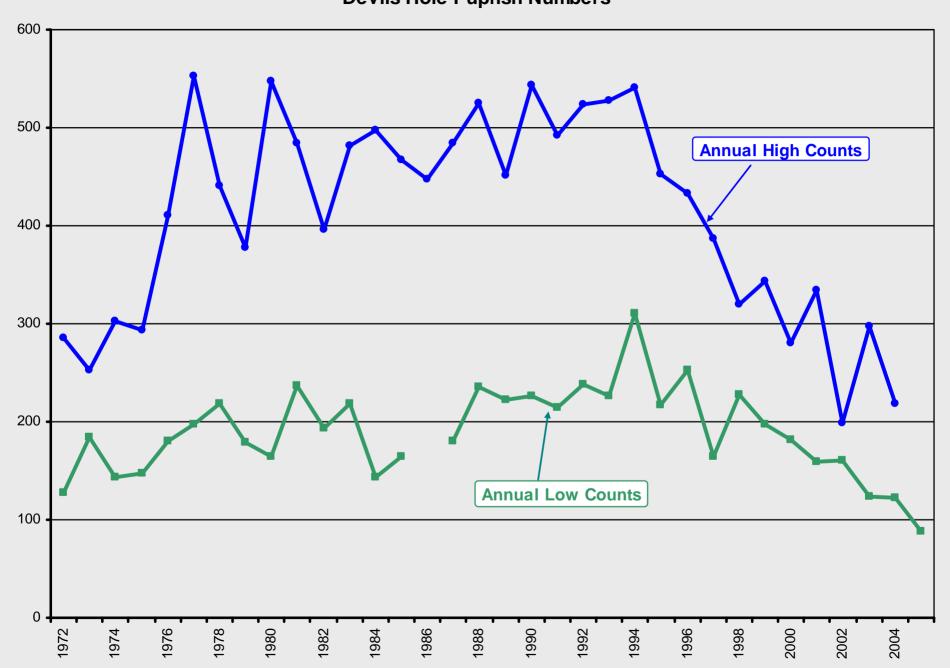




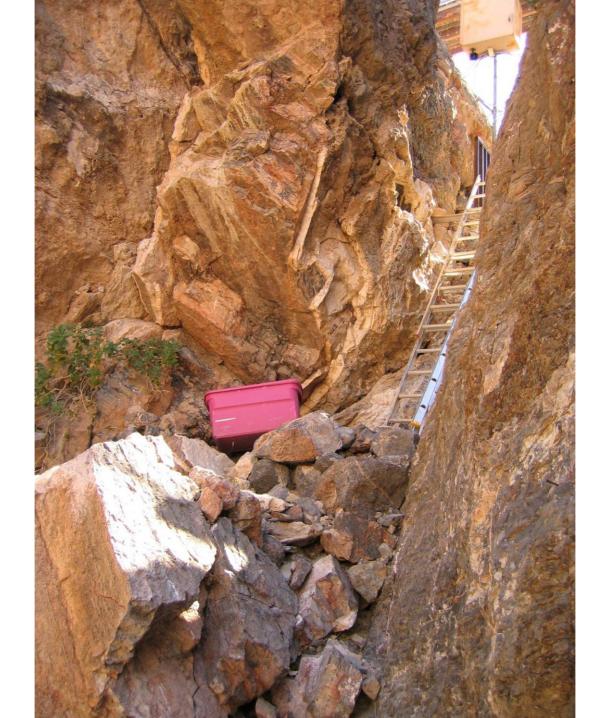




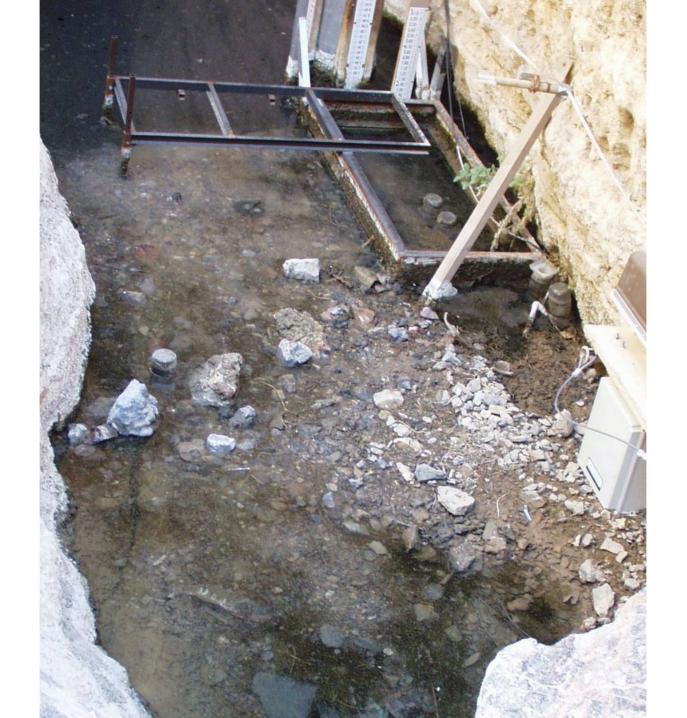
Devils Hole Pupfish Numbers

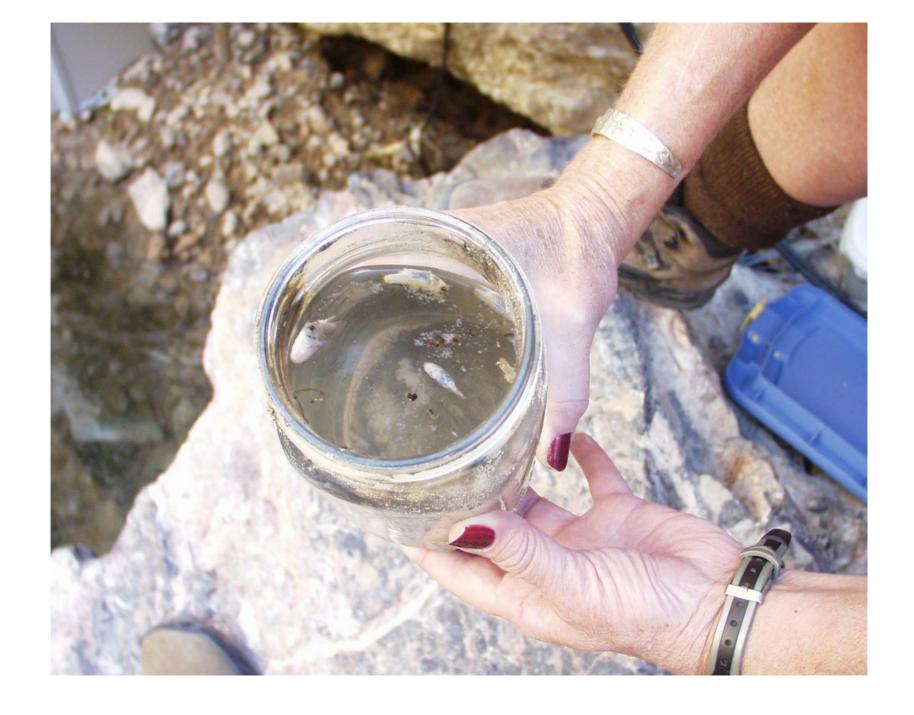


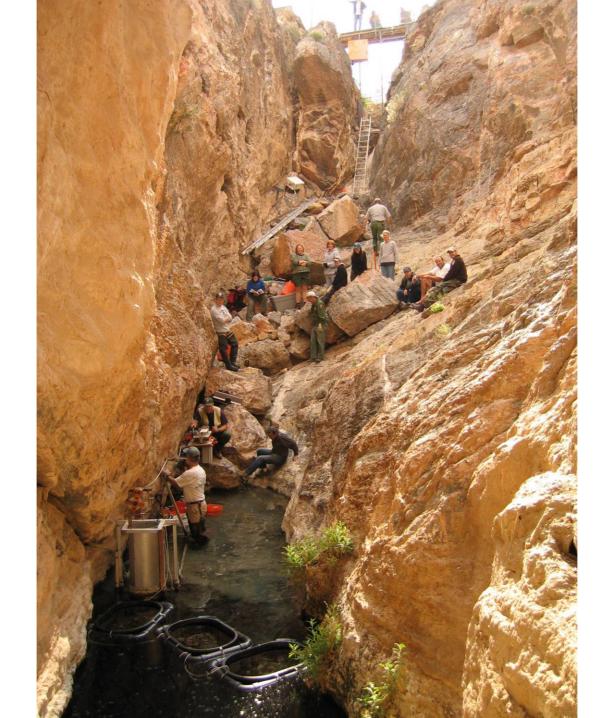














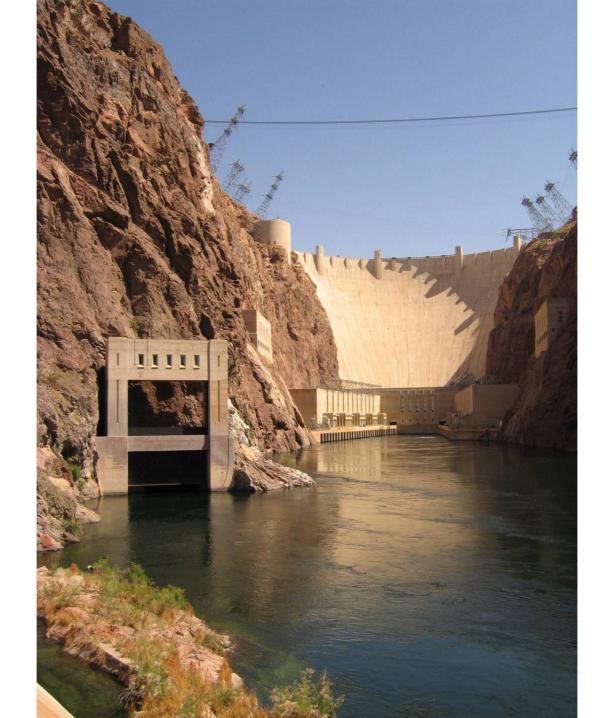
















Devils Hole Pupfish Populations

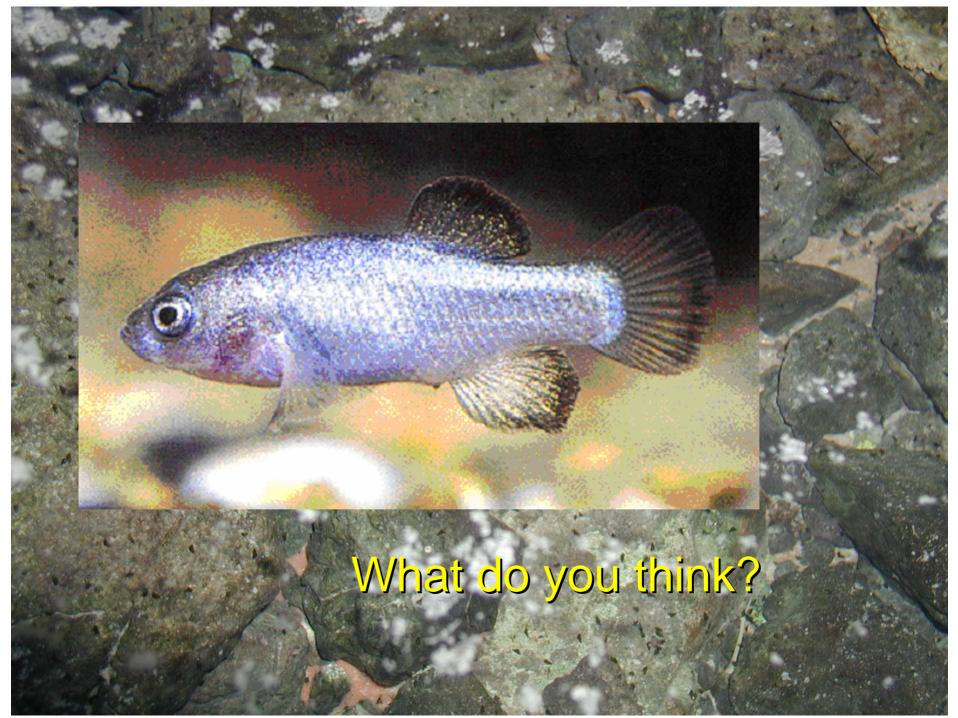
Devils Hole (DH) Death Valley NP	84 mature
Point of Rocks (PoR) FWS Refuge	180 (likely hybrids with <i>C. nevadensis</i>)
Hoover Dam (HD) BOR Refuge	50 mature

Estimates of population genetic differentiation using R_{ST} above the diagonal and F_{ST} below the diagonal

	DH	HD	PoR '98	PoR '05
DH		0.108*	0	0.593***
HD	0.072***		0.032	0.365***
PoR	0.007	0.044**		0.513***
1998				
PoR	0.179**	0.221**	0.182**	
2005				

Possible Recovery Actions

- Remove some Devils Hole fish to Hoover Dam and expand population.
- 2. Remove some Devils Hole fish to Point of Rocks, and backcross with current stock.
- 3. Add fish to Devils Hole from either refuge.
- 4. Add no fish and monitor Devils Hole.





Captive Propagation Program for the Key Largo Woodrat (Neotoma floridana smalli)







- One of 5 named subspecies of the Eastern woodrat
- Classified as a federally endangered species in 1984 due to concerns over habitat loss and the impact of commercial development
- Nocturnal herbivore, feeding on the buds, leaves, and fruit of many plant species
- "Females can have up to two litters a year, consisting of one to four young with an average of two"

Subspecies Justification

160 miles

Virtually no hardwood hammock between localities

Morphological differences

Molecular taxonomy unknown





Population Decline

- •During the late 70's and early 80's stick nests were abundant in North Key Largo
- •In 1986 Humphrey estimated the KLWR population to be ~6,500 individuals.
- •Large decline appeared to occur during the late 80's and early 90's when there was no trapping occurring





Potential Threats:



Habitat Loss and degradation

Habitat management

Disease

Introduced predators and competitors

Interactions of factors and demographic effects



Habitat Loss and Fragmentation }

More then 47% of woodrat habitat has been lost to development, today Key Largo woodrats are restricted to approximately 850 hectares of protected tropical hardwood hammock forest in northern Key Largo

Rock Harbor

Previous Range of the Key Largo Woodrat

Card

Rd.

Ocean

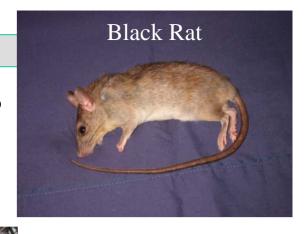
Reef Club



Predators and Competitors



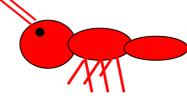
Yellow Rat Snake /











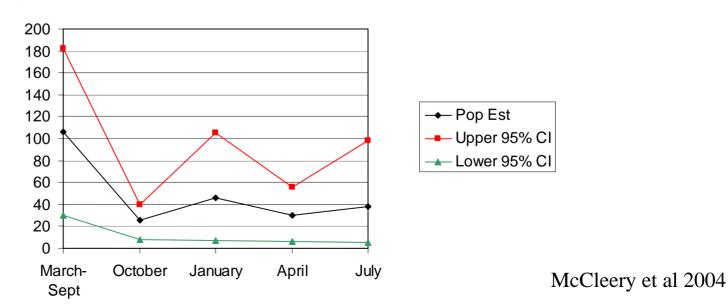




Current Population Estimates

By 1995 stick nests had all but disappeared from North Key Largo

The KLWR population was estimated at 106 (30-182) in 2002, an October 2004 report estimated the current population at 40 (5-104).

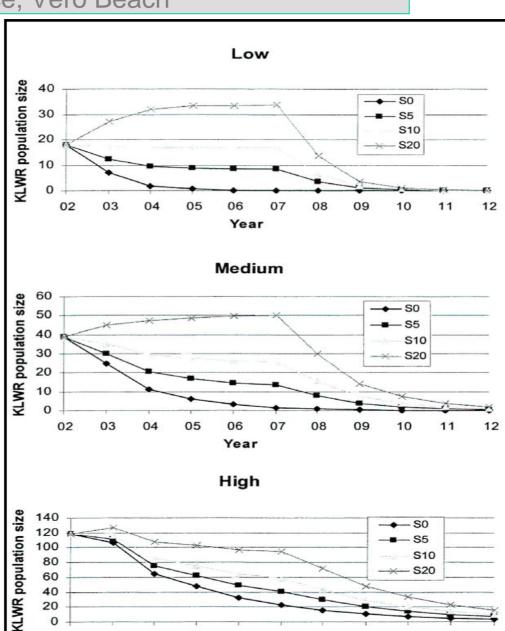




2003 PVA (McCleery et al 2004)

Predicts a 70 to 98 % risk of extinction in 10 years regardless of "best case scenario" or "worst case scenario" parameters

Introduction of 20 females annually for 5 years may prevent extinction, although once introductions are stopped the PVA continues to predict extinction



Year

08

10

02

12

In April 2002, we initiated an experimental captive propagation program and began developing a captive propagation and reintroduction plan in accordance with Service policy.



HERBERT



SOPHIE



Captive Propagation and Reintroduction Plan

Objectives:

Prevent the extinction of KLWRs in the wild by maintaining the population through augmentation with captive-born woodrats

Maintain present levels of genetic diversity (i.e. fitness) and heterozygosity

Captive Propagation and Reintroduction Plan

Objectives:

Maximize the conservation of genetic diversity in both the wild and captive populations by developing appropriate measures to safeguard and increase diversity

Maintain refugia populations in captivity until causes of the current decline are understood and addressed

Maintain refugia populations in captivity in case of catastrophic events



After proving husbandry techniques we brought two more woodrats into captivity and began attempting to breed woodrats.





BILL FELICIA



Woodrats are territorial and will fight to the point of causing serious injury, therefore all woodrats are housed separately and all breeding attempts are

observed by zoo staff







SUCCESS!!

5/10/03 Sophie and Bill became the proud parents of two female pups

5/23/03 Felicia and Bert became the proud parents of a single male pup

ONLY after demonstrating breeding success did the RO authorize us to fully implement the Plan and bring in 12 woodrats from the wild







With breeding success came the need for

Genetics-based Captive Breeding Management

In order to meet the objectives of the Captive Propagation and Reintroduction Plan all woodrats are genotyped at microsatellite DNA loci to establish relatedness to other captive individuals. Pairwise genetic relatedness (based on multilocus genotypes) determines potential matings. The ultimate objective is to avoid inbreeding depression while maximizing genetic diversity and heterozygosity.



Genetic Analysis

- Closest known relative is the Eastern woodrat (Neotoma floridana)
- This non-listed subspecies served as a "surrogate species" for training Service and Commission biologists to take blood samples and insert microchip identification tags
- No phylogenetic analyses comparing the two subspecies have been reported (research currently underway)
- Compare genetic diversity within and between subspecies to place observed levels of *N. f. smalli* genetic diversity into proper phylogenetic perspective.

Methods

Trapping performed by USFWS Ecological Services

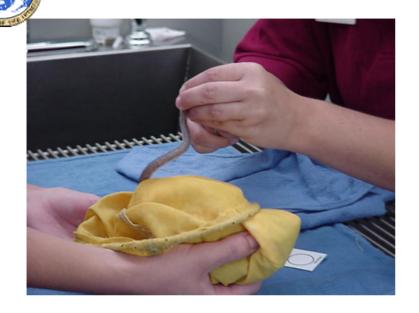
Blood taken and preserved on FTA cards; tail tips also placed in ethanol

Fecal samples placed on FTA cards and in alcohol

Genotyping performed by USGS

Microsatellite DNA markers developed for congeneric species (Allegheny woodrat)

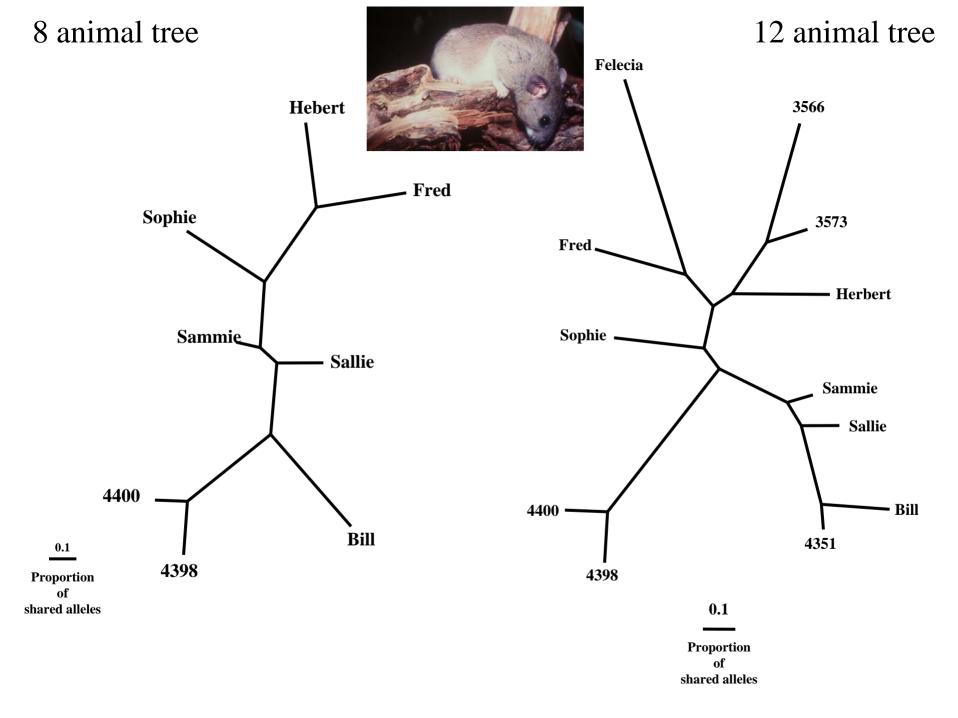
Nine markers currently in use; others being developed





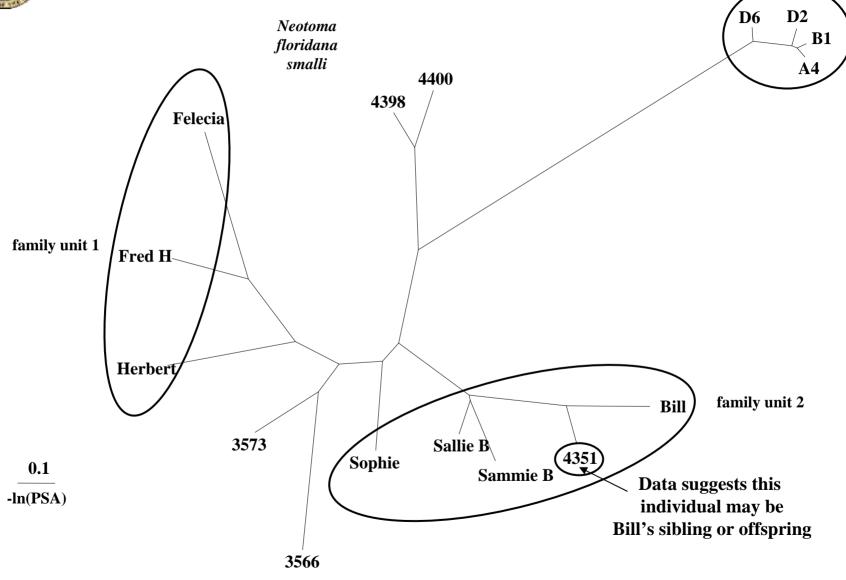








Neotoma floridana floridana



Unrooted neighbor-joining tree

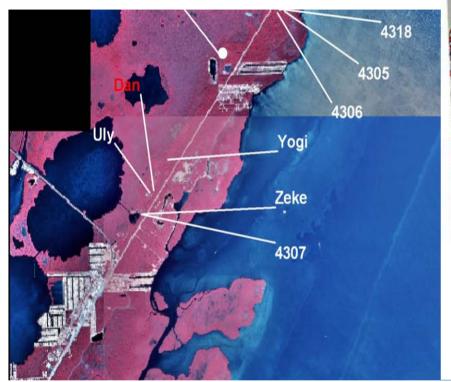
Captive Neotoma floridana smalli February 24, 2004 Fred H Family 1a Felic Family 1b -Herbert - deceased (father of both Frannie Fred H and Frannie) **Sophie** Sallie 1 Nick Sammie B Ripper Family 2 Family 3 Rita Bill 0.1 **Females**

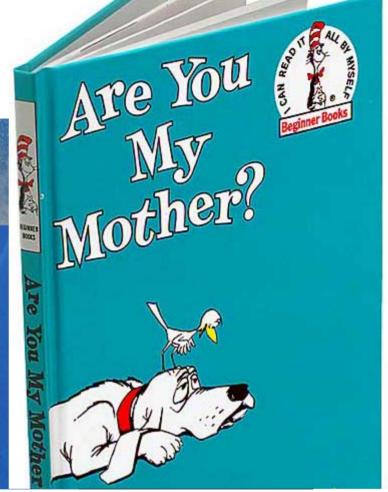
PSA Distance units

Males

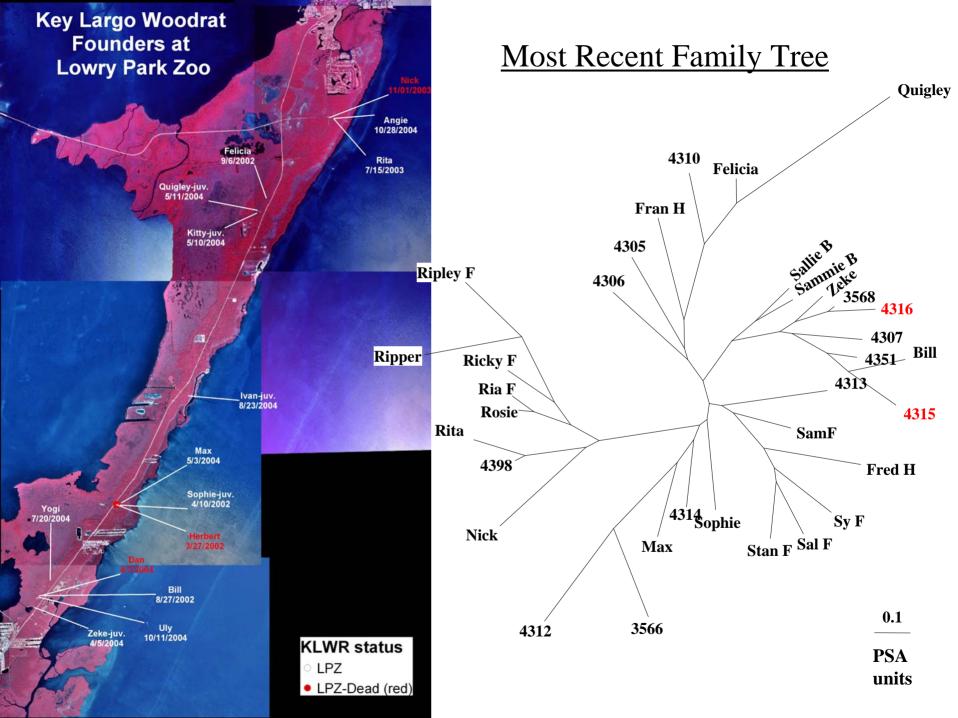
Animal 1	Animal 2	<u>PSA</u>	Relationship								
Molly	Jamie	0.251	siblings			Scenario A					
Ripper	Nick	0.492	unknown	Site 1	Site 2		Site 1	PSA		Site 2	PSA
Rita	Ripper	0.492	mother-daughter	Nick	Bill	Sophie	Nick	1.099	Bill	Rita	0.811
Bert	Sam	0.588	father-son	Sam	Bert	Molly	Nick	1.281	Bill	Felecia	1.792
Bert	Sophie	0.588	related?	Jamie	Rita	Jamie	Nick	1.504	Bill	Ripper	1.281
Bill	Jamie	0.588	father-daughter	Molly	Felecia	Sophie	Sam	0.693	Bert	Felecia	1.099
Bill	Molly	0.588	father-daughter	Sophie	Ripper	Sam	Jamie	1.099	Bert	Rita	1.281
Rita	Nick	0.588	related?		[Fran]	Sam	Molly	1.281	Bert	Ripper	1.504
Sam	Felecia	0.588	mother-son				Mean	1.160		Mean	1.295
Sophie	Jamie	0.588	mother-daughter				Std	0.273		Std	0.336
Sophie	Molly	0.588	mother-daughter								
Sophie	Sam	0.693	related?								
Bill	Rita	0.811	related?			Scenario B					
Bert	Jamie	0.944	unknown	Site 1	Site 2		Site 1	PSA		Site 2	PSA
Bert	Molly	0.944	unknown	Nick	Bert	Felecia	Nick	1.792	Bert	Rita	1.281
Bill	Nick	0.944	unknown	Bill	Sam	Sophie	Nick	1.099	Bert	Ripper	1.504
Molly	Felecia	0.944	unknown	Felecia	Rita	Bill	Felecia	1.792	Bert	Jamie	0.944
Sophie	Felecia	0.944	unknown	Sophie	Ripper	Bill	Sophie	1.281	Bert	Molly	0.944
Sophie	Rita	0.944	unknown	[Fran]	Jamie		Mean	1.491	Sam	Rita	1.281
Bert	Felecia	1.099	paired family 1		Molly		Std	0.355	Sam	Ripper	1.504
Bert	Nick	1.099	unknown						Sam	Jamie	1.099
Sam	Jamie	1.099	unknown						Sam	Molly	1.281
Sophie	Nick	1.099	unknown							Mean	1.230
Sophie	Ripper	1.099	unknown							Std	0.220
Bert	Rita	1.281	unknown								
Bill	Ripper	1.281	unknown								
Bill	Sophie	1.281	paired family 2			Scenario C					
Jamie	Felecia	1.281	unknown	Site 1	Site 2	The second secon	Site 1	PSA		Site 2	PSA
Jamie	Rita	1.281	unknown	Nick	Bill	Molly	Nick	1.281	Bill	Rita	0.811
Molly	Nick	1.281	unknown	Bert	Sam	Jamie	Nick	1.504	Bill	Ripper	1.281
Molly	Rita	1.281	unknown	Molly	Rita	Felecia	Nick	1.792	Bill	Sophie	1.281
Sam	Molly	1.281	unknown	Jamie	Ripper	Bert	Molly	0.944	Sam	Rita	1.281
Sam	Nick	1.281	unknown	Felecia	Sophie	Bert	Jamie	0.944	Sam	Ripper	1.504
Sam	Rita	1.281	unknown		[Fran]	Bert	Felecia	1.099	Sophie	Sam	0.693
Bert	Ripper	1.504	unknown				Mean	1.261		Mean	1.142
Bill	Bert	1.504	unknown				Std	0.337		Std	0.316
Jamie	Nick	1.504	unknown								
Molly	Ripper	1.504	unknown								
Sam	Ripper	1.504	unknown								
Bill	Felecia	1.792	unknown								
Bill	Sam	1.792	unknown								
Felecia	Nick	1.792	unknown								
Felecia	Ripper	1.792	unknown								
Jamie	Ripper	1.792	unknown								
Rita	Felecia	1.792	unknown								

Parentage Question





W oodrat	N m a 1	N m a 1	N m a 4	Nma42	N m a 5	N m a 5	Nm a8	Nm a8	Nma10	Nma10	Nma11	Nm a 11	Nm a 14	Nm a14	Nm aD21	Nm aD21	N m a D 138	Nm aD138	
4307	321	321	165	171	231	243	132	138	213	225	164	166	161	163	155	155	293	297	Mom - ?
Zeke	315	321	159	165	231	231	132	138	213	225	164	164	161	161	151	155	293	297	
Dan	301	321	153	159	243	243	134	138	209	225	162	164	163	167	151	155	293	293	Dad - no





Controlled Propagation Prospects and Obstacles

Successful Breeding – 14 pups, 6 single pup births and 4 "twin" births; captive-born individuals have been successfully mated both to each other and to wild-caught individuals

Genetic analysis – high level of diversity

Have brought in all 12 wild-caught individuals allowed by the Plan (Total 13 male, 13 female)

KLWRs will be split into 2 captive populations, endocrinology lab





Controlled Propagation Prospects and Obstacles

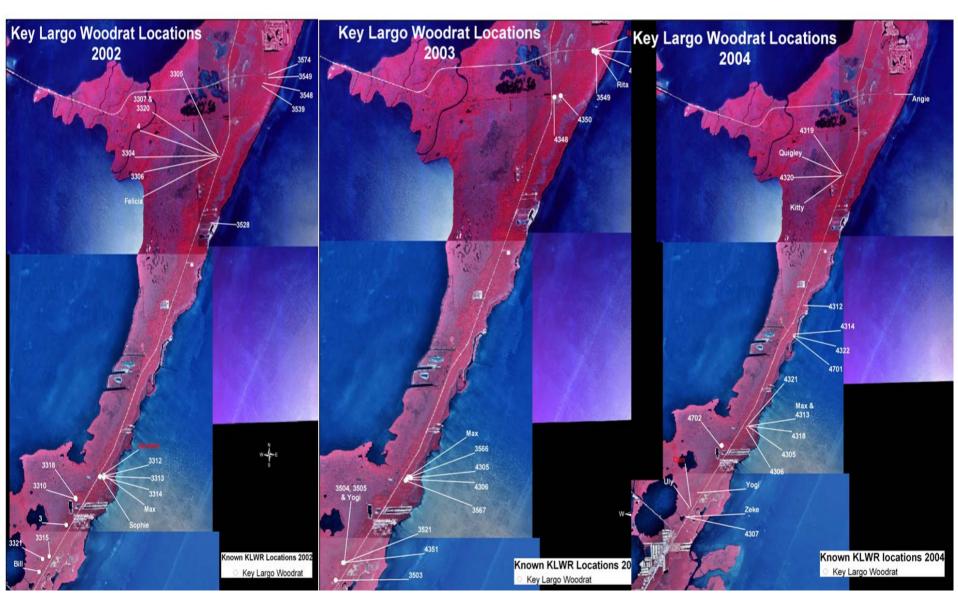
Captive aggression increasing – loss of one wild-caught male

Causes of decline yet to be fully identified or addressed

Hesitant to release captive animals until we understand and address the causes and issues behind the decline of the wild population

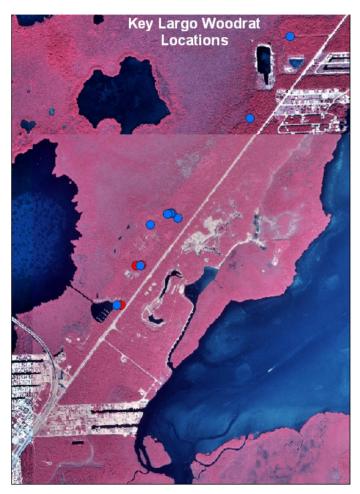


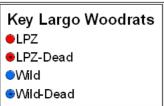


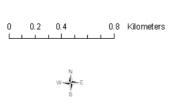




Woodrat vs. Black Rat Locations 2002-2004

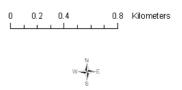














Recent Success in the Wild

2002 - 13 "new" woodrats captured

2003 - 14 "new" woodrats captured

2004 – 20 "new" woodrats captured, 10 of which were captured in October and November

First stick nest identified last month from an area where more then 30

raccoons have been removed!







Partners:

USFWS

USGS

Florida Fish and Wildlife Conservation Commission

Florida Department of Environmental Protection

Lowry Park Zoo

Texas A&M University

University of Georgia

Copyrighted zoological facility outside of Orlando